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SEQUENCE LISTING
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<110> The Chemo-Sero-Therapeutic Research Institute

<120> A DNA coding for merozoite protein of Babesia cabally, a recombinant protein

obtained by using said DNA and a use thereof

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<211> 1828

<212> DNA

<213> Babesia caballi

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344

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Val Gly Asp Val Thr Lys Thr Leu Leu Ala Ser Glu Ser Val Asp

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Ser Ala Ala Asn Ala Tyr Met Ile Asn Ser Asp Met Ser Asp Tyr Leu

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TCG GCT GTG TCT GAC AAC TTC GCQ GAG CGC ATT TGC AGT CAG GTC CCT

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25

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Lys Gly Ser Asn Cys Ser Ma Ser Val Ser Ala Tyr Met Ser Arg Cys

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65 70

GCT AAA CAG GAC TGC CTG ACT CTC CAA AGT CTT AAG TAC CCT CTT GAG

Ala Lys Gln Asp Cys Veu Thr Leu Gln Ser Leu Lys Tyr Pro Leu Glu

85

GCT AAG TAC CAA COG CTG ACC CTT CCT GAC CCC TAC CAG TTG GAG GCC

Ala Lys Tyr Gln ₱ro Leu Thr Leu Pro Asp Pro Tyr Gln Leu Glu Ala

95

GCA TTT ATA CTC TTC AAG GAG AGT GAC GCT AAT CCG GCC AAT AGC ACT

Ala Phe Ile Leu Phe Lys Glu Ser Asp Ala Asn Pro Ala Asn Ser Thr

110

105

115

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GAG AAG CGC TTC TGG ATG CGT TTC AGA AGG GGC AAG AAC CAC AGT TAC Glu Lys Arg Phe Trp Met Arg Phe Arg Arg Gly Lys Asn His Ser Tyr TTC CAC GAC TTA GTC TTC AAT CTG CTG GAG AAG AAC GTG AQT CGC GAC Phe His Asp Leu Val Phe Asn Leu Leu Glu Lys Asn Val Ahr Arg Asp GCG GAT GCT ACT GAC ATT GAG AAC TTT GCG TCC AGG TAC CTG TAC ATG Ala Asp Ala Thr Asp Ile Glu Asn Phe Ala Ser Arg/Tyr Leu Tyr Met GCC ACG CTT TAC TAC AAG ACG TAC ACG AAT GTT GAT GAG TTC GGT GCT Ala Thr Leu Tyr Tyr Lys Thr Tyr Thr Asn Val/Asp Glu Phe Gly Ala AGC TTC TTT AAC AAG TTG TCT TTC ACT ACT CGG TTG TTC GGC TGG GGC Ser Phe Phe Asn Lys Leu Ser Phe Thr Thy Gly Leu Phe Gly Trp Gly ATC AAG AGG GCA CTT AAG CAG ATT ATT/CGC TCT AAC CTG CCC CTT GAC Ile Lys Arg Ala Leu Lys Gln Ile Ile Arg Ser Asn Leu Pro Leu Asp ATC GGG ACA GAA CAC AGC GTC AGT/ CGC CTG CAG CAC ATT ACG AGC AGT Ile Gly Thr Glu His Ser Val Ser Arg Leu Gln His Ile Thr Ser Ser TAC AAG GAT TAC ATG GAT ACE CAG ATT CCT GCA CTG CCC AAG TTT GCG Tyr Lys Asp Tyr Met Asp Thr Gln Ile Pro Ala Leu Pro Lys Phe Ala AAG CGT TTC TCC CTT ATG GTA GTG CAG AGG CTG CTG GCC ACC GTG GCT Lys Arg Phe Ser Leu Met Val Val Gln Arg Leu Leu Ala Thr Val Ala GGT TAC GTC GAC ACC CCG TGG TAT AAG AAG TGG TAC ATG AAG CTG AAG Gly Tyr Val Asp/Thr Pro Trp Tyr Lys Lys Trp Tyr Met Lys Leu Lys AAC TTT ATG OTG AAC AGG GTG TTC ATT CCT ACA AAG AAG TTC TTC AAT Asn Phe Met/Val Asn Arg Val Phe Ile Pro Thr Lys Lys Phe Phe Asn AAG GAA ATT CGT GAG CCT AGT AAG GCA TTA AAA GAA AAG GTG TCA ACC Lys Gly Ile Arg Glu Pro Ser Lys Ala Leu Lys Glu Lys Val Ser Thr

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Phe Phe Asn Lys Glu Ile Arg Asp Pro Ser Lys Ala Leu Lys/Glu Lys	
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215

Gln His Ile Thr Ser Ser Tyr Lys Asp Tyr Met Asp Thr Gln Ile Pro

220

1828

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